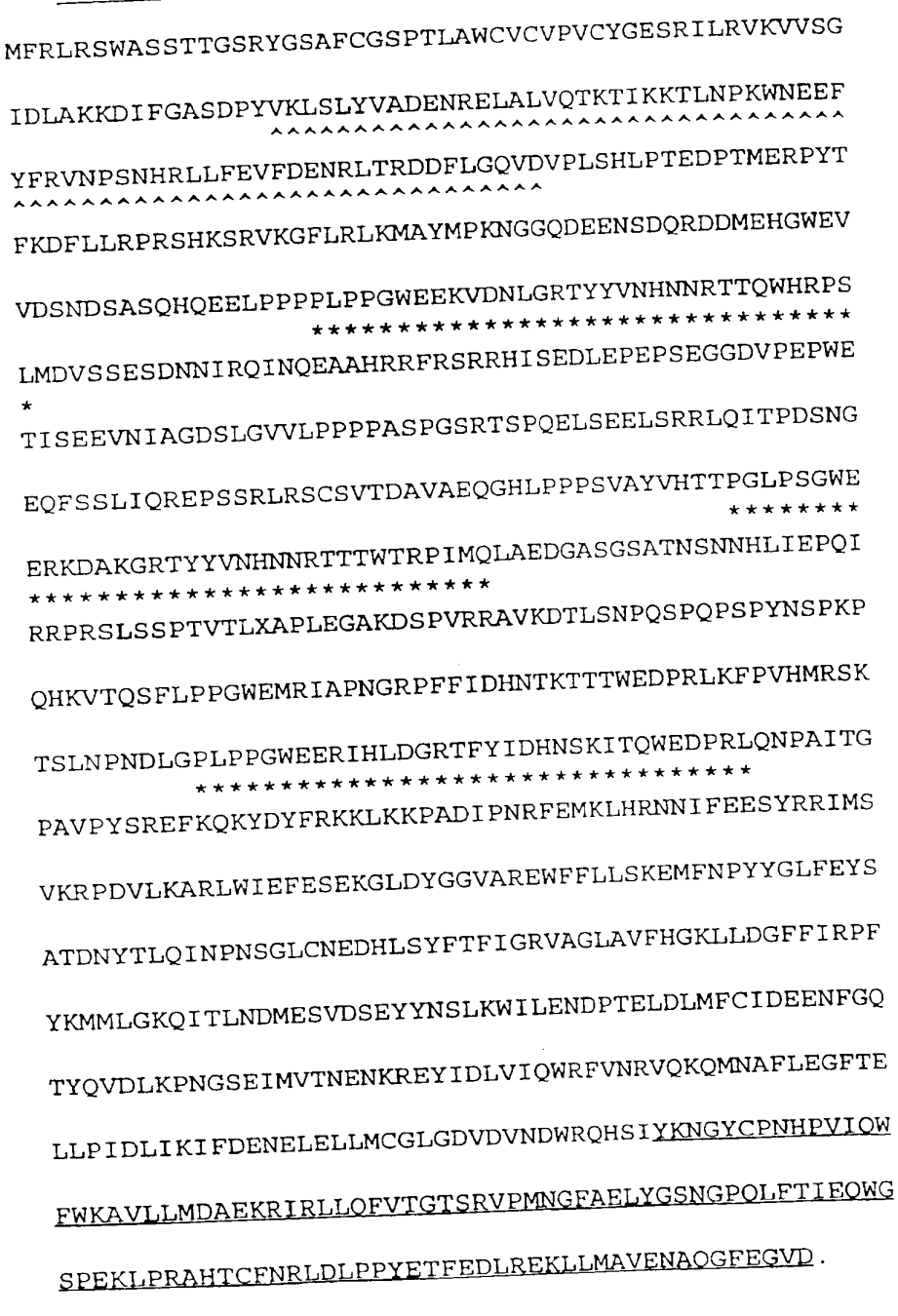
FIGURE 1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FKDFLLRPRSHKSRVKGFLRLKMAYMPKNGGQDEENSDQRDDMEHGWEV
VDSNDSASQHQEELPPPPLPPGWEEKVDNLGRTYYVNHNNRTTQWHRPS
LMDVSSESDNNIRQINQEAAHRRFRSRRHISEDLEPEPSEGGDVPEPWE
TISEEVNIAGDSLGVVLPPPPASPGSRTSPQELSEELSRRLQITPDSNG
EQFSSLIQREPSSRLRSCSVTDAVAEQGHLPPPSVAYVHTTPGLPSGWE
ERKDAKGRTYYVNHNNRTTTWTRPIMQLAEDGASGSATNSNNHLIEPQI
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QHKVTQSFLPPGWEMRIAPNGRPFFIDHNTKTTTWEDPRLKFPVHMRSK
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PAVPYSREFKQKYDYFRKKLKKPADIPNRFEMKLHRNNIFEESYRRIMS
VKRPDVLKARLWIEFESEKGLDYGGVAREWFFLLSKEMFNPYYGLFEYS
ATDNYTLQINPNSGLCNEDHLSYFTFIGRVAGLAVFHGKLLDGFFIRPF
YKMMLGKQITLNDMESVDSEYYNSLKWILENDPTELDLMFCIDEENFGQ
TYQVDLKPNGSEIMVTNENKREYIDLVIQWRFVNRVQKQMNAFLEGFTE
LLPIDLIKIFDENELELLMCGLGDVDVNDWRQHSIYKNGYCPNHPVIQW
FWKAVLLMDAEKRIRLLQFVTGTSRVPMNGFAELYGSNGPQLFTIEQWG
SPEKLPRAHTCFNRLDLPPYETFEDLREKLLMAVENAQGFEGVD.

FIGURE 2

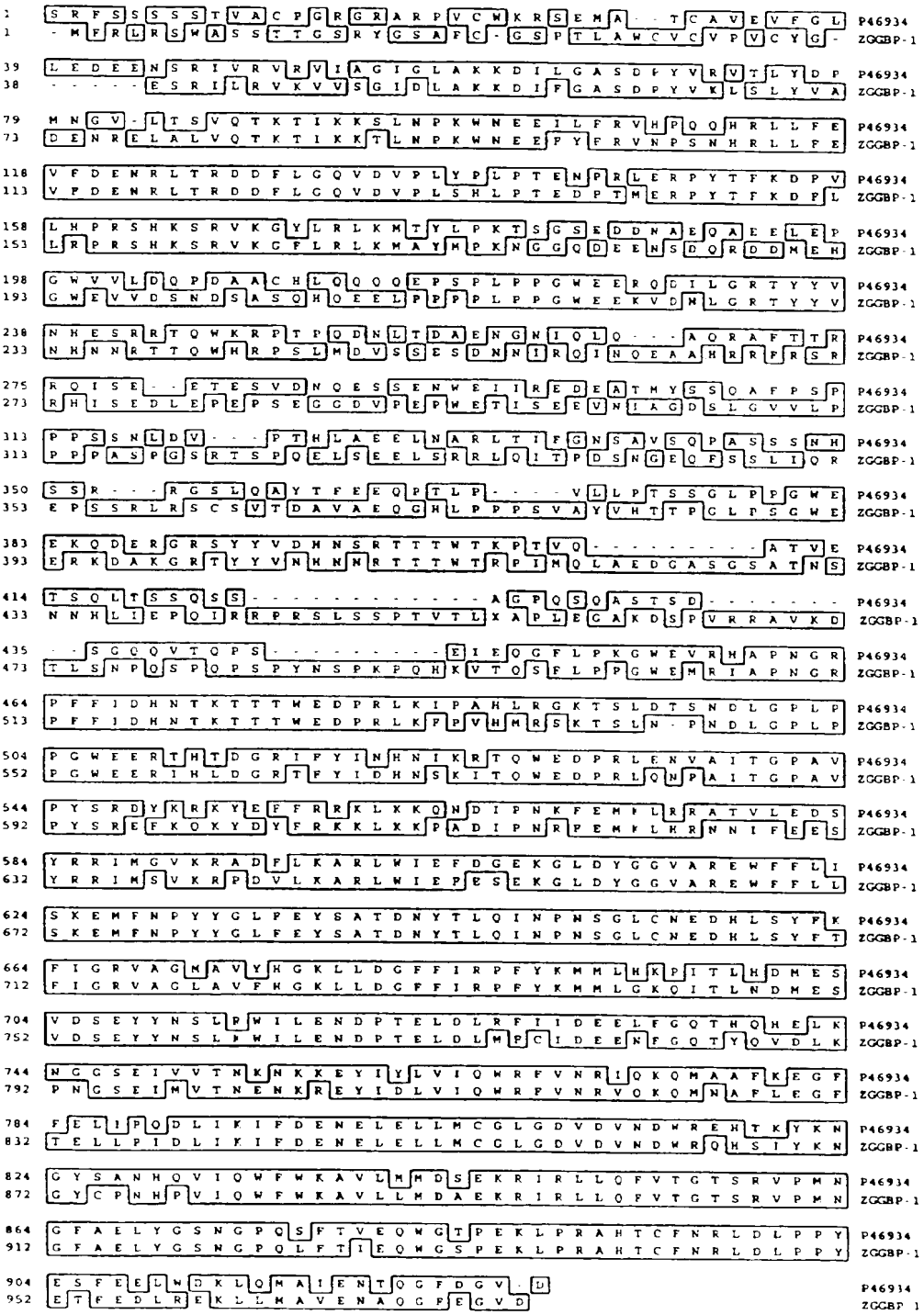


FIGURE 3

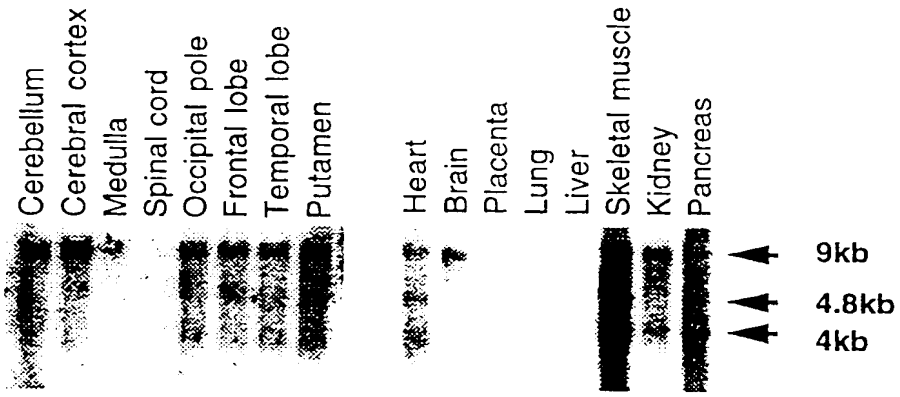

Cerebellum
Cerebral cortex
Medulla
Spinal cord
Occipital pole
Frontal lobe
Temporal lobe
Putamen
Heart
Brain
Placenta
Lung
Liver
Skeletal muscle
Kidney
Pancreas
9kb
4.8kb
4kb

FIGURE 4

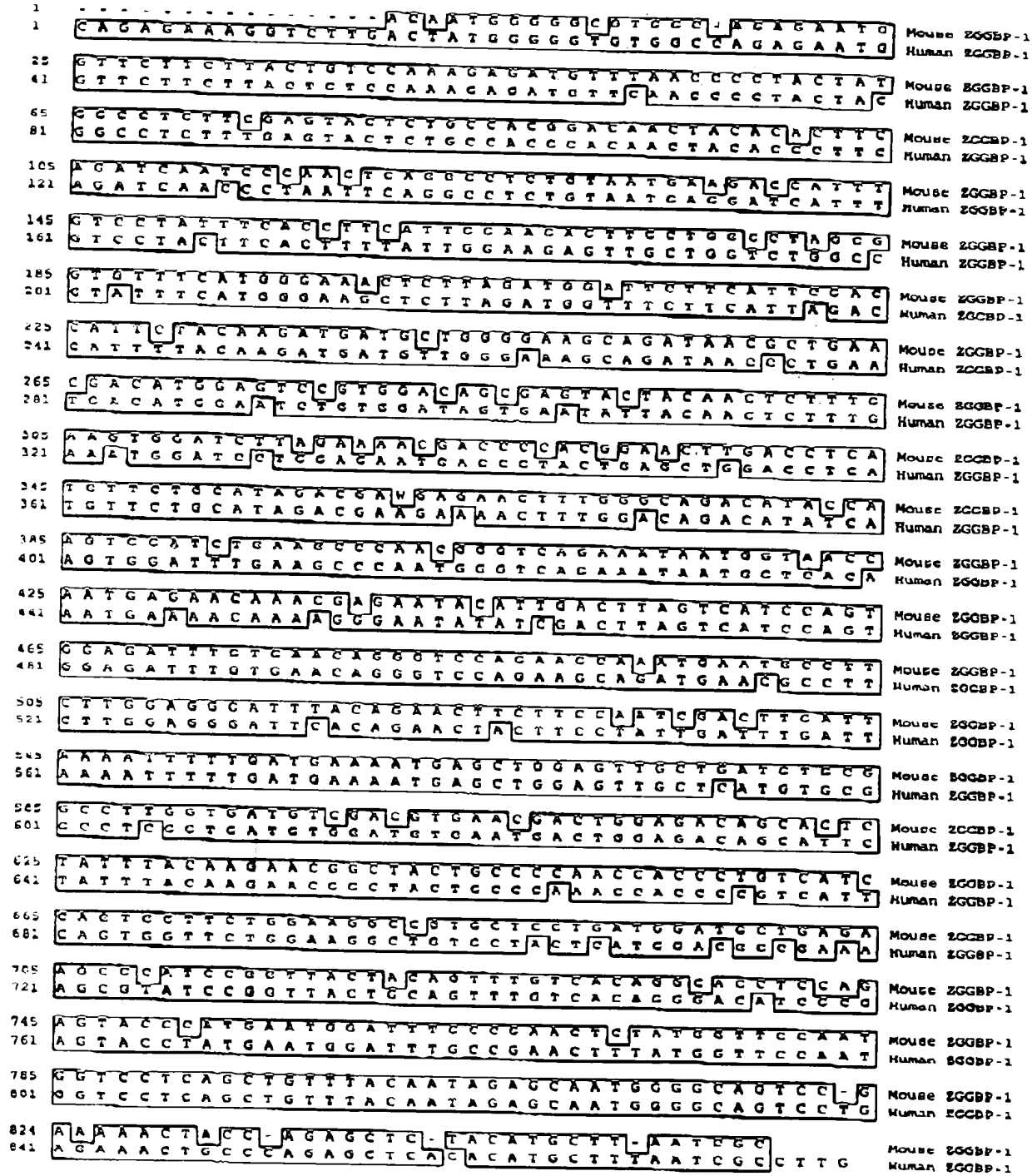
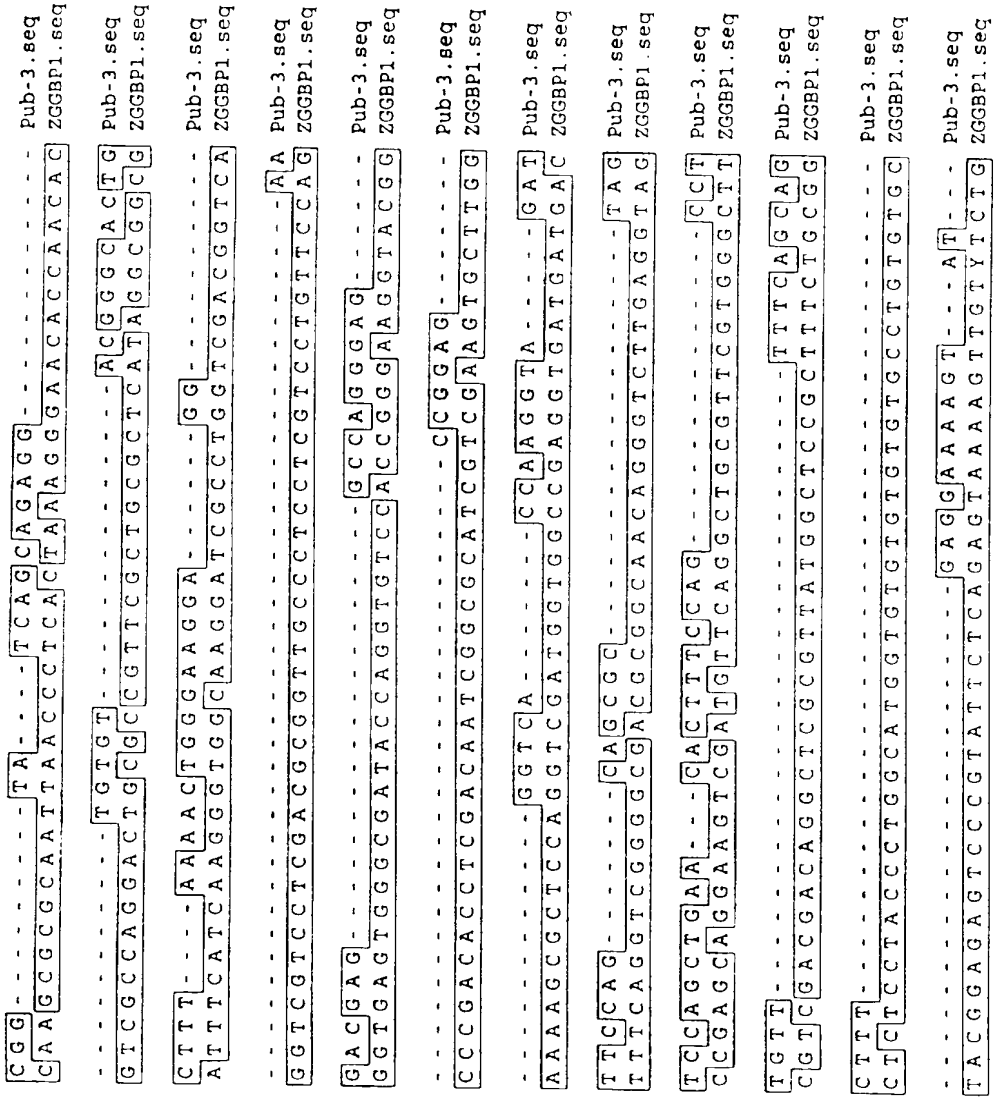


FIGURE 5



09/463844

FIGURE 5 continued

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521  TCCGTATGTGAAACTTTCATTGTACGTAGCGGATGAGAAT  ZGGBP1.seq

202  AGAGAACTTGCTTTGGTCCAGACAAAAACAATTAAAAAGA  Pub-3.seq
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242  CACTGAACCCAAAATGGAATGAAGAATTTTATTTCAGGGT  Pub-3.seq
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282  AAACCCATCTAATCACAGACTCCTATTTGAAGTATTTGAC  Pub-3.seq
641  AAACCCATCTAATCACAGACTCCTATTTGAAGTATTTGAC  ZGGBP1.seq

322  GAAAATAGACTGACACGAGACGGCTTCCTGGGCCAGGTGG  Pub-3.seq
681  GAAAATAGACTGACACGAGACGACTTCCTGGGCCAGGTGG  ZGGBP1.seq

362  ACGTGCCCCTTAGTCACCTTCCGACAGAAGATCCAACCAT  Pub-3.seq
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FIGURE 5 continued

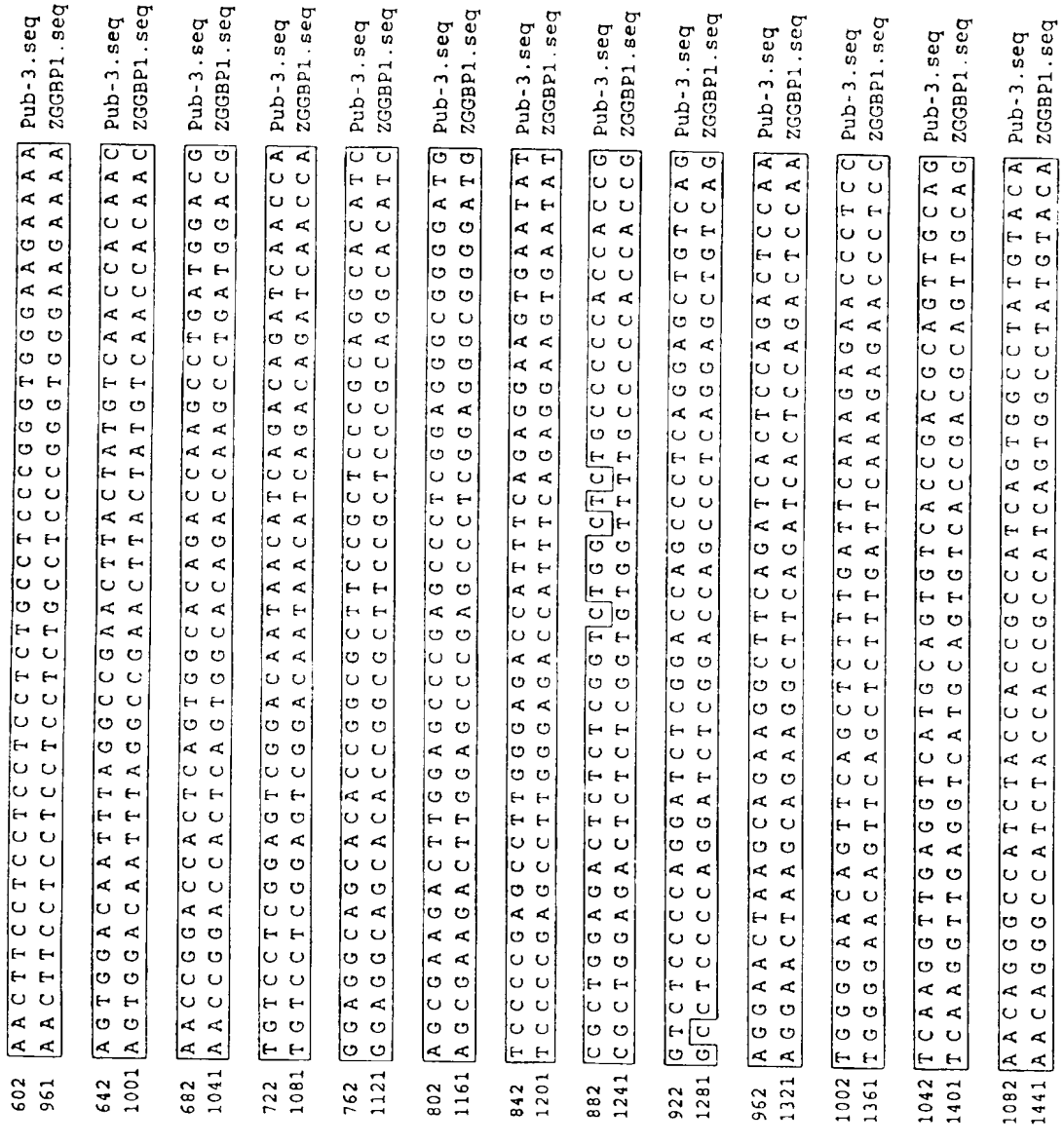

FIGURE 5 continued

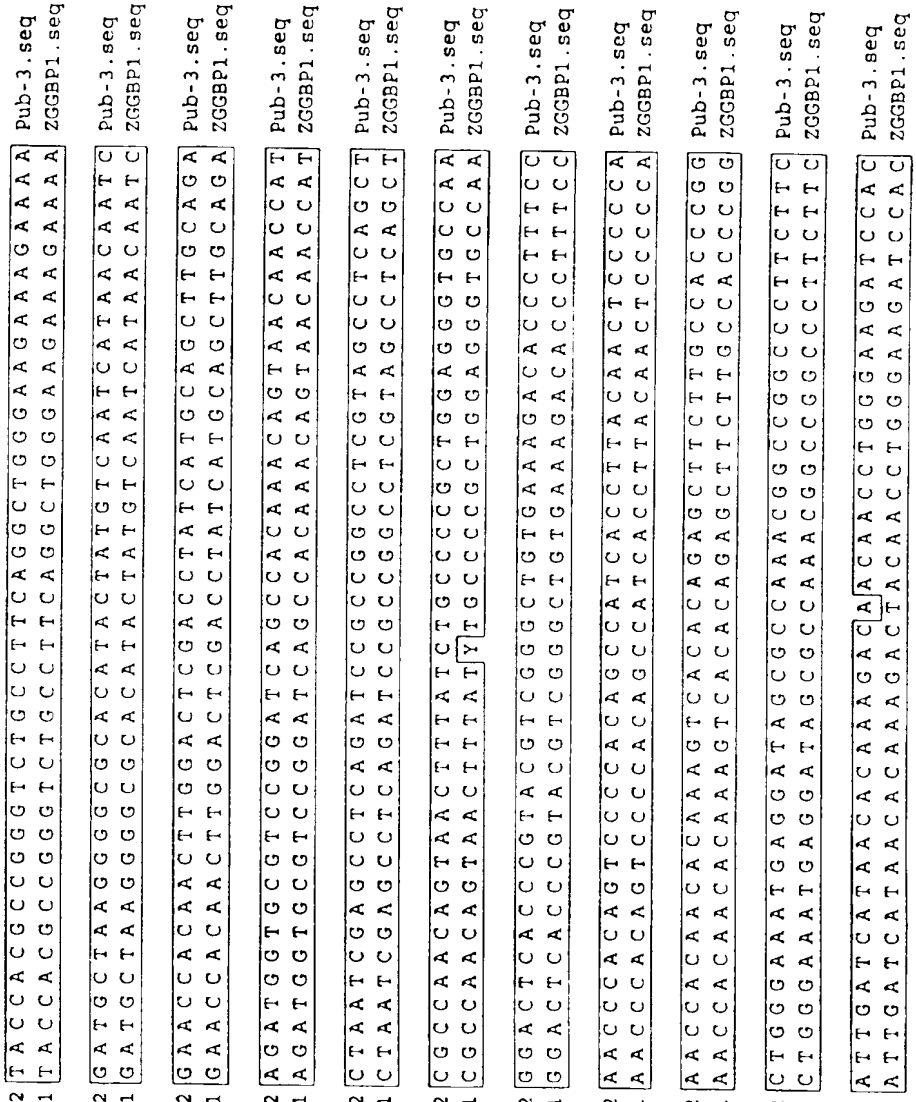

FIGURE 5 continued

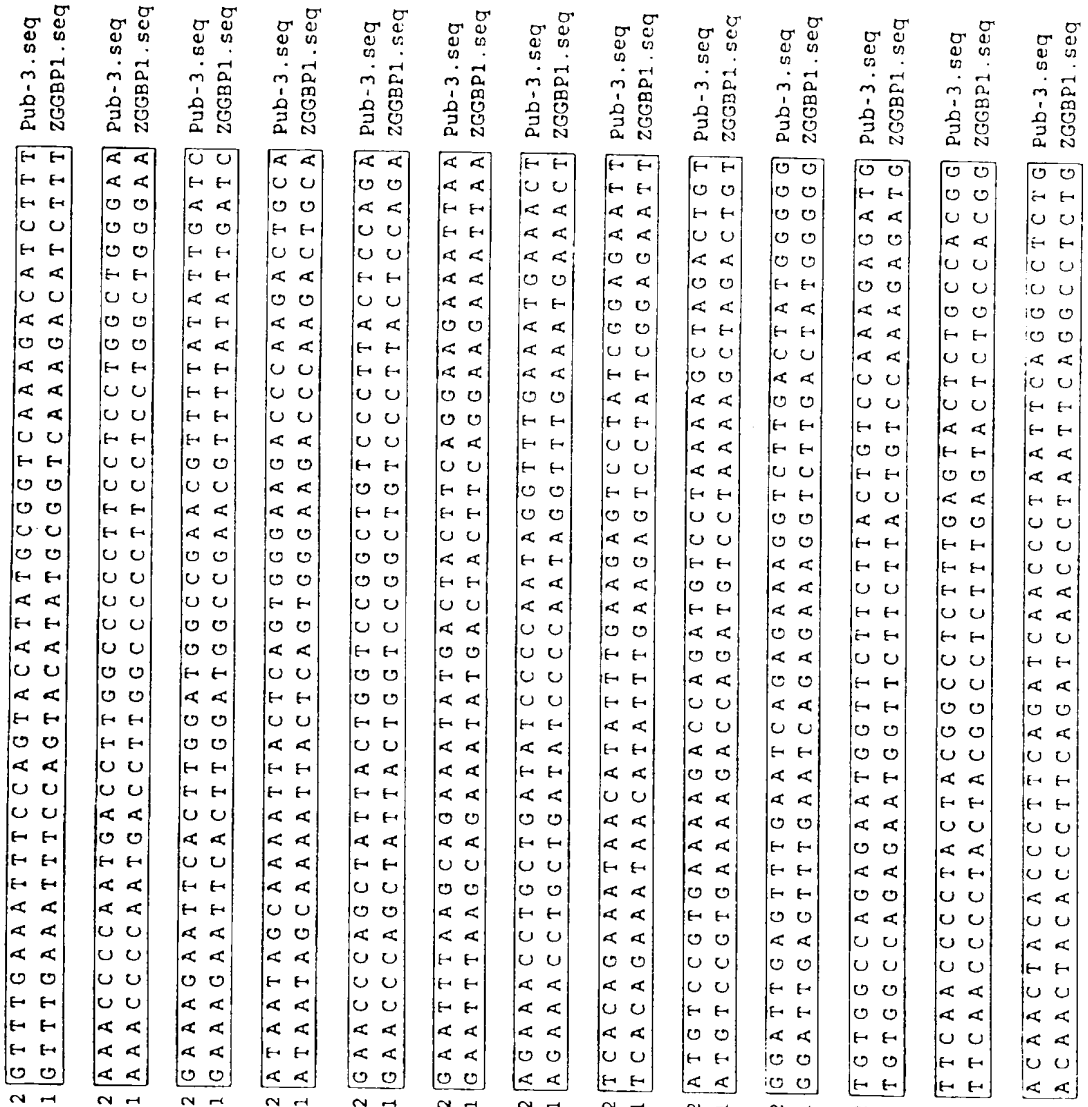

FIGURE 5 continued

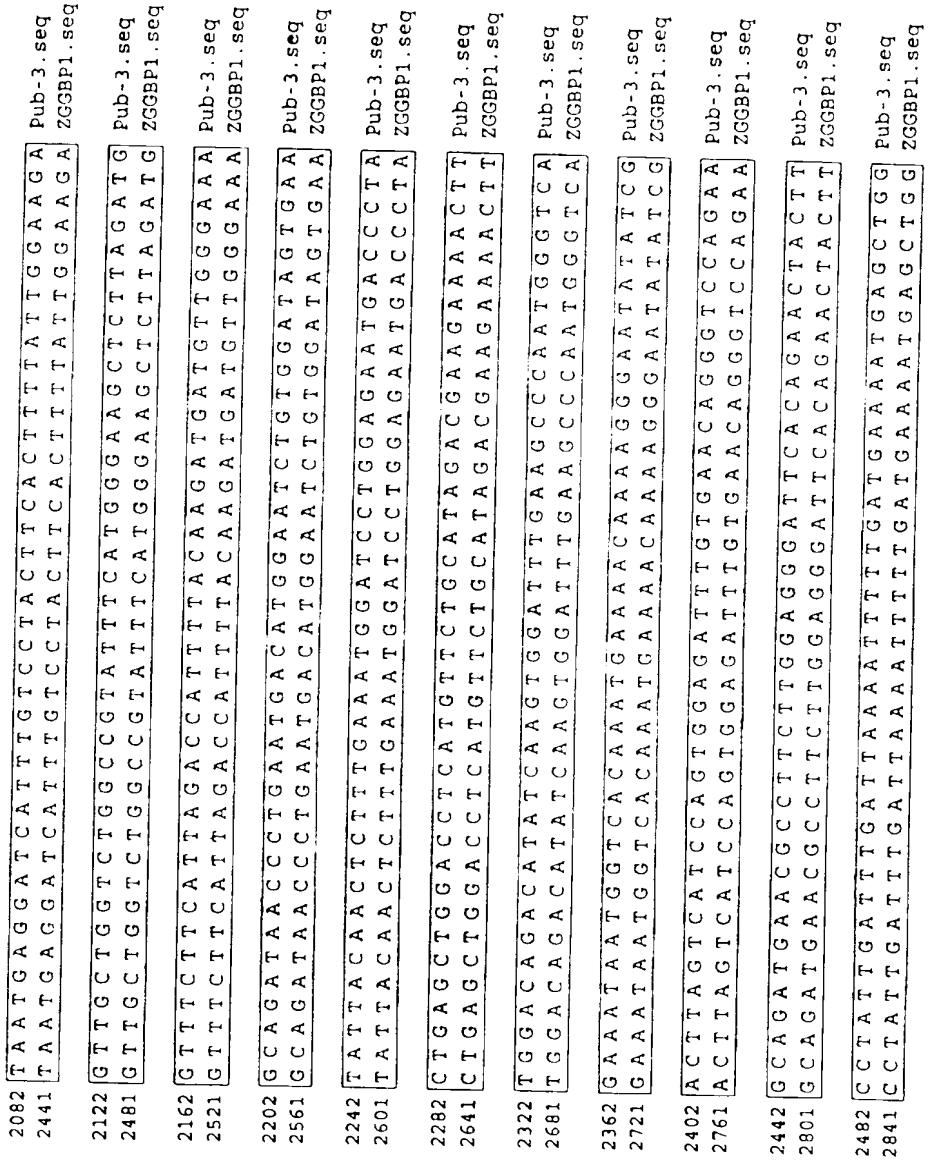

FIGURE 5 continued

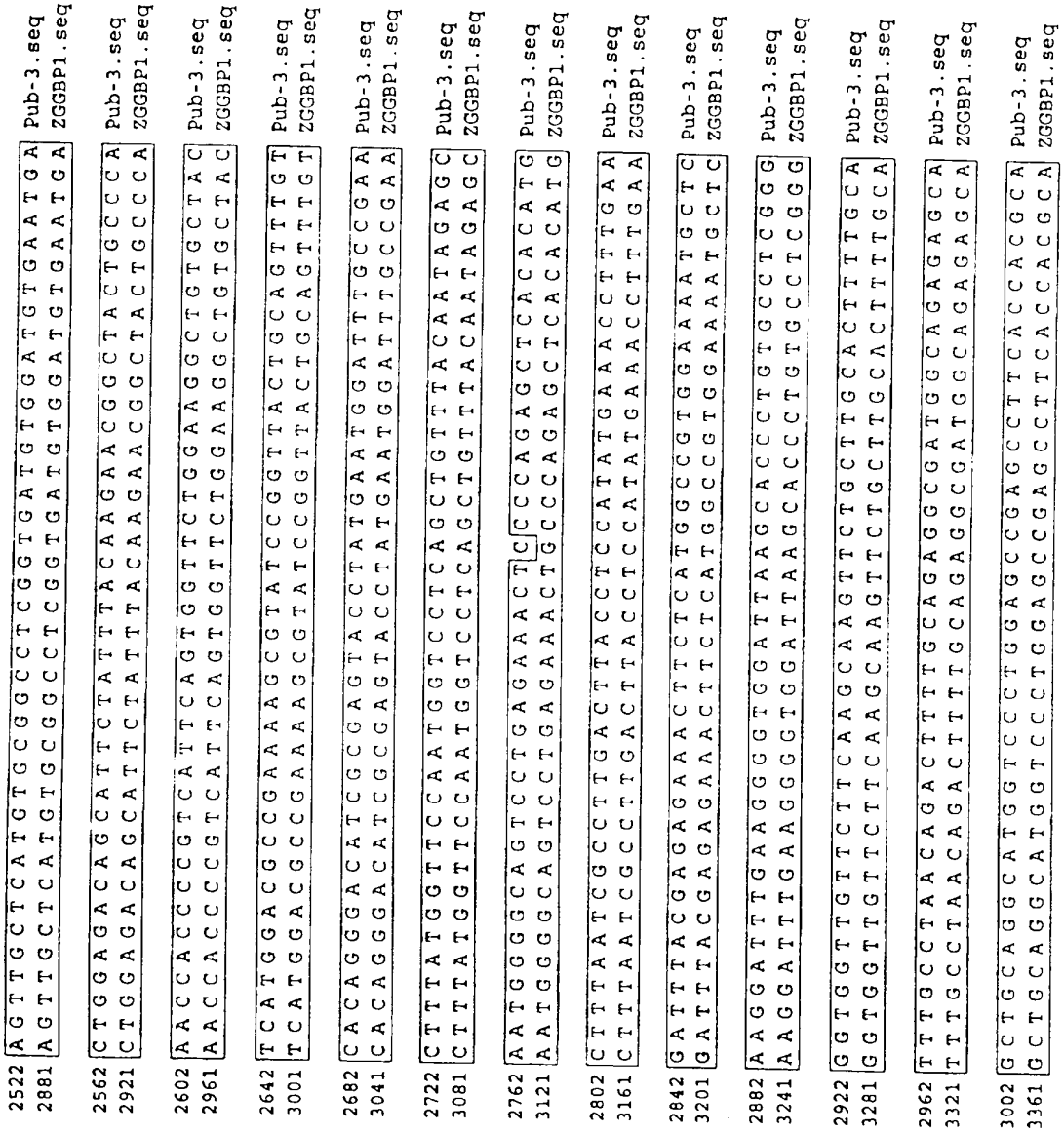

FIGURE 5 continued

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3214 ---------------------------------------- Pub-3.seq
3960 CTAATAGCTACAGGCTGAGAGAATTGTAACATAGCATGAC ZGGBP1.seq

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3214 ---------------------------------------- Pub-3.seq
4040 TTAGAAGTAATTACATGTGTTCTAACACATTTGAGACAGG ZGGBP1.seq

3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
4120 TCCTGGGCGCTGTACAGTCATCTTTTATTCTATTTCCTCT ZGGBP1.seq

3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
4240 TCCTGTTTGTATGCGTTTGCTAATGGTAGCTAAATAACCA ZGGBP1.seq

3214 ---------------------------------------- Pub-3.seq
4280 GTTTTTGTTGTAAATGCACCAATTCTGAAGGCACTTTATG ZGGBP1.seq

3214 ---------------------------------------- Pub-3.seq
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FIGURE 5 continued

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4440 GGCTGCTGGTGTATAAAAACCTGGATGTAAAGCTGAGCCT ZGGBP1.seq

3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
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3214 ---------------------------------------- Pub-3.seq
4760 TCGTGCGTTTGAGAAAACACCTGAGGCAGTGTGGGAGTTG ZGGBP1.seq
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FIGURE 5 continued

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3214 ---------------------------------------- Pub-3.seq
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3226                                          Pub-3.seq
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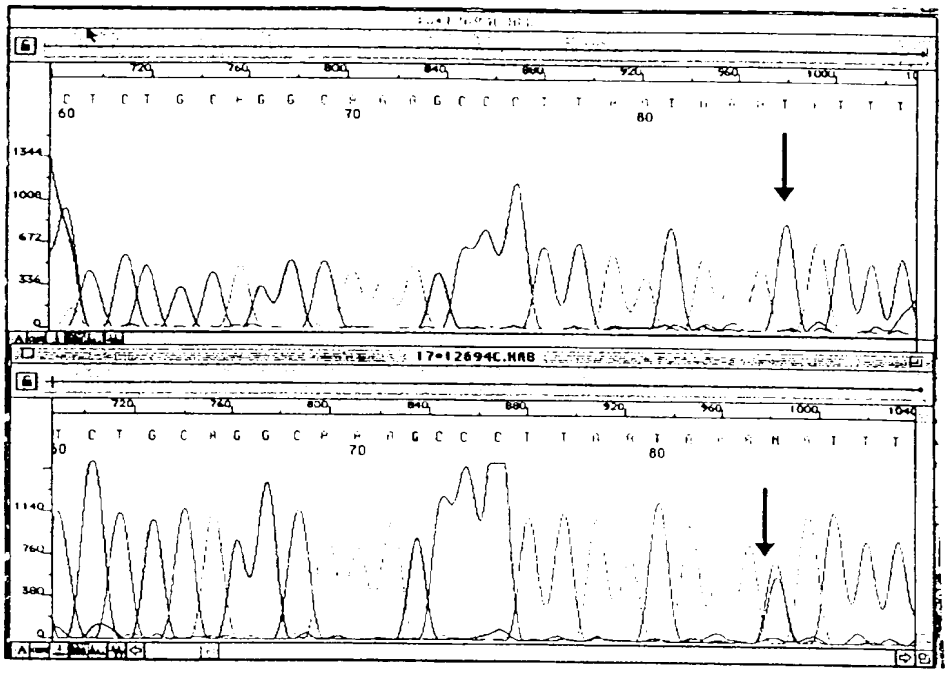
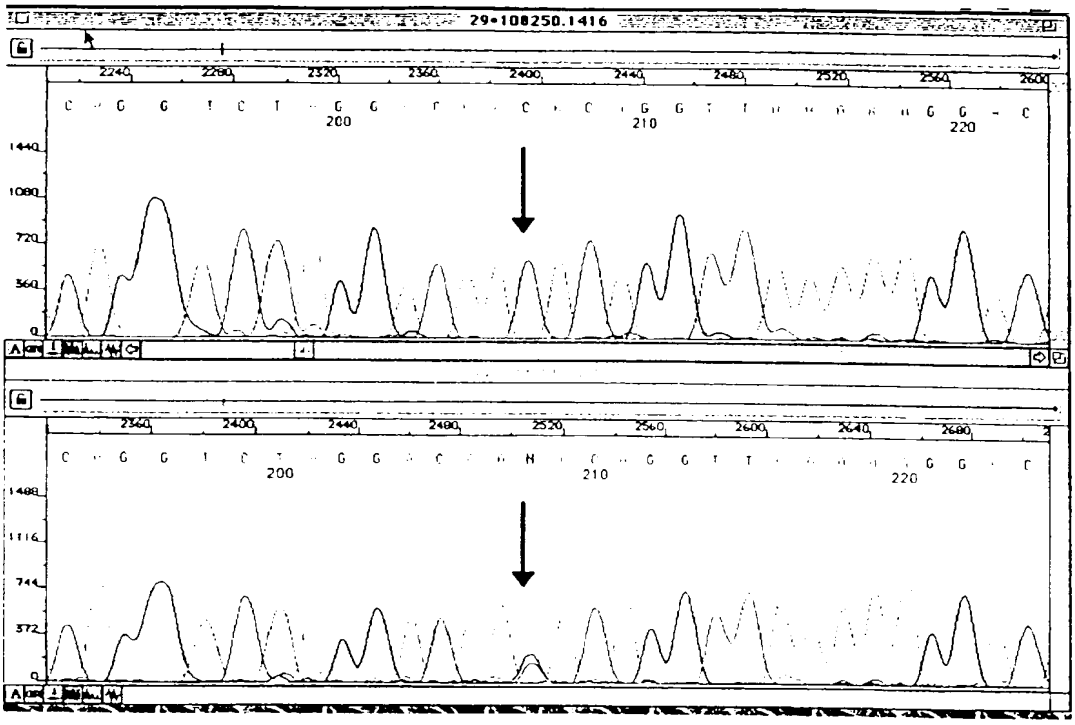
Decoration 'Decoration #1': Box residues that match the Consensus exactly.



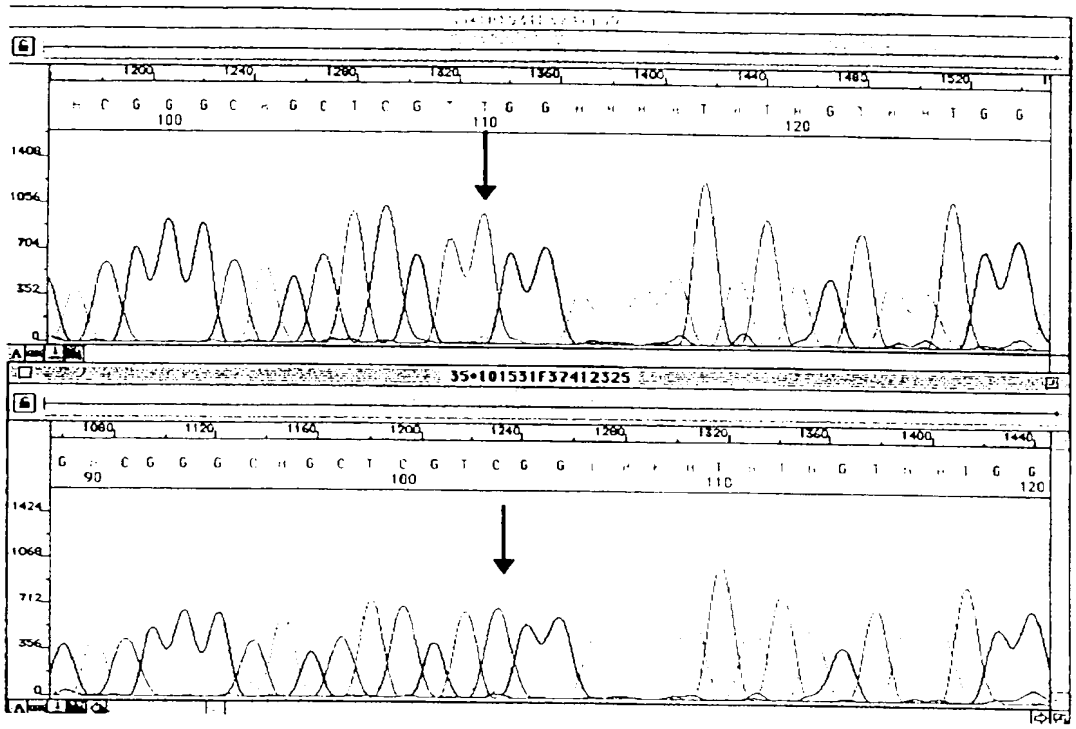
Wild Type (human foetal brain)	**T/T**
Variant Type (human adult brain)	**T/C**
Polymorphism Position	**3554**
RFLP	**-**

FIGURE 6



Wild Type (GM1416)	C/C
Variant (7225)	C/G
Position	4828

FIGURE 7



Primer sequences derived from BAC and used on lymphoblastoid cell lines from BPAD Patients.

Homozygous wild type (KK169) - T/T

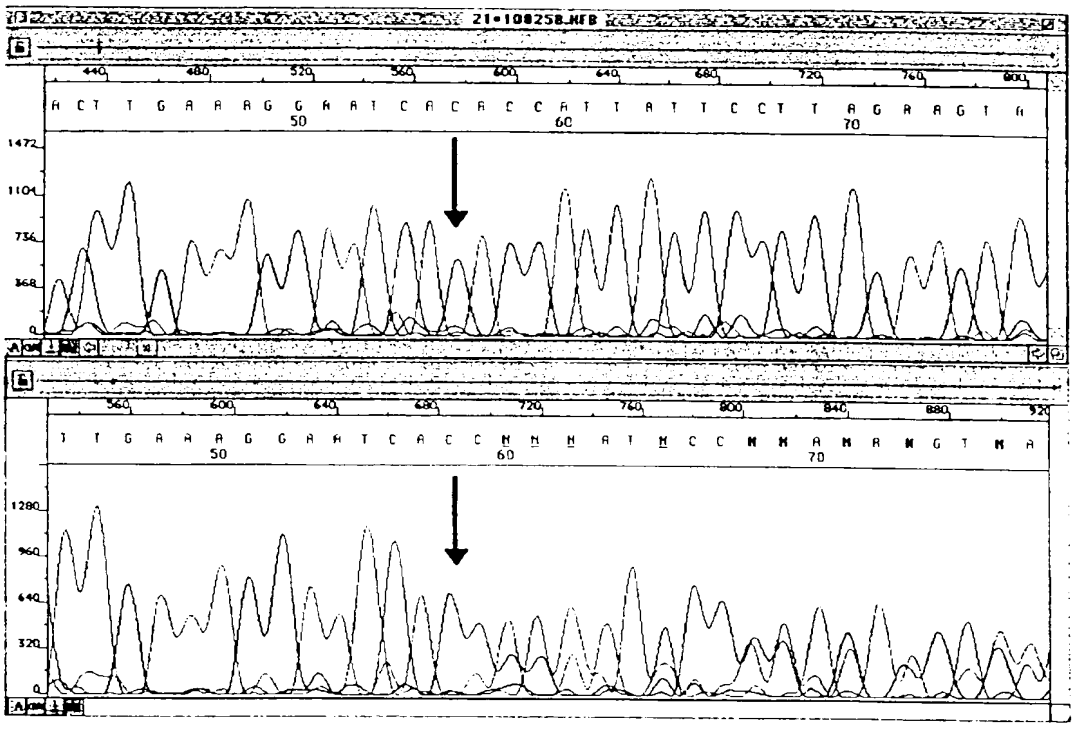
Homozygous variant (KK232) - C/C

FIGURE 8

Figure 9

TGCTGCAAGTGACAGGTTCCAAGAAGCCCGAGGGCTCAGAGCTGAATGATGAAGCGC
AGTCCCCAAAGTGCCTGGCCACCCCTCCCTCCCTGGATCACTGCTGCCTGGGCTTGA
TTGATTGATTGATTGATTGATTGATTTTGAGAGAGATTCTCACTGTCACCCAG
GCTGGAGTACAGTGGTGCGATCTCGGCTCACTGCAGCCTCTGCCTCCCGGGTTCAAG
CAATTCTCCTGCCTCAGCCTCCCAAGTAGCTGGGACTACAGGCACGCGCCACCACAC
CCAGCTAATTTTGTATTTTTAGTAAAAGACGGGGTTTCACCATGTTGGGCCAGGATG
GTCTTGATCTCCTGACCTCATGATCCACCCGCCCCGGCTTCCAAAGTGCTGGGATAC
AGGCATGAACCCGACGCGCCCAGCATGGACATTTTTTTTTAATCCCCTGCCCTTTTC
TTGNGGCATAATTCATTGCAGGTCTCTTCTATACAGATCATGGAAAACACATTTTCT
TAACTGAGTTNTTATTATTTATACCCAGNCACCTCATGACANNTTTACCCTGTTACA
NACAAAATGGGCACCTGCCAAAANCAACTTTNATATAAGGATGCTCCAGGCCT

Tetranucleotide repeat underlined



Top electropherogram (human foetal brain) - wild type

Lower electropherogram (7225) - heterozygous variant

Arrow indicates the position of the C+C insertion - position 4032

FIGURE 10